SEQUENCE LISTING

<110> Takeshi TABIRA Hideo HARA

5

<120> A Recombinant Adeno-Associated Virus Vector for Treatment of Alzheimer Disease

<130> 40072-0026US

10

<140> US 10/560,280

<141> 2005-12-12

<150> PCT/JP2004/008224

15 <151> 2005-06-11

<150> JP 2003-169714

<151> 2003-06-13

20 <150> JP 2003-371103

<151> 2003-10-30

<160> 19

25 <170> PatentIn Ver. 2.0

<210> 1

<211> 129

<212> DNA

30 <213> Homo sapiens

<220>

<221> CDS

<222> (1)..(129)

<400> 1 gat gca gaa ttc cga cat gac tca gga tat gaa gtt cat cat caa aaa Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys 1 10 15 5 5 ttg gtg ttc ttt gca gaa gat gtg ggt tca aac aaa ggt gca atc att Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile 20 25 30 gga ctc atg gtg ggc ggt gtt gtc ata gcg aca 129 10 Gly Leu Met Val Gly Gly Val Val Ile Ala Thr 40 35 <210> 2 15 <211> 43 <212> PRT <213> Homo sapiens 20 <400> 2 Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys 1 5 10 15 Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile 25 20 25 30 Gly Leu Met Val Gly Gly Val Val Ile Ala Thr 35 40 30 <210> 3 <211>63 <212> DNA <213> Homo sapiens 35

<220> <221> CDS <222> (1)..(63) 5 <400> 3 gat gca gaa ttc cga cat gac tca gga tat gaa gtt cat cat caa aaa Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys 1 5 10 15 ttg gtg ttc ttt gca 63 10 Leu Val Phe Phe Ala 20 <210> 4 15 <211> 21 <212> PRT <213> Homo sapiens <400> 4 20 Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys 1 5 10 15 Leu Val Phe Phe Ala 25 20 <210> 5 <211> 54 30 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(54) 35

```
<400> 5
     atg ctg ccc ggt ttg gca ctg ctc ctg ctg gcc gcc tgg acg gct cgg
     Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala Trp Thr Ala Arg
                                       10
                                                          15
       1
                      5
 5
                                                                 54
     gcg ctt
     Ala Leu
10
     <210> 6
     <211> 18
     <212> PRT
     <213> Homo sapiens
     <400> 6
15
     Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala Trp Thr Ala Arg
                      5
                                       10
                                                          15
       1
     Ala Leu
20
     <210> 7
     <211> 197
25
     <212> DNA
     <213> Artificial Sequence
     <220>
     <223> Description of Artificial Sequence: DNA Construct
30
     <220>
     <221> CDS
     <222> (9)..(191)
35
     <220>
     <221> misc_feature
```

```
<222> (3)..(8)
     <223> Xba I recognition site
     <220>
 5 <221> sig_peptide
     <222> (9)..(62)
     <400> 7
     ggtctaga atg ctg ccc ggt ttg gca ctg ctc ctg ctg gcc gcc tgg acg 50
10
              Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala Trp Thr
                1
                               5
                                                  10
     gct cgg gcg ctt gat gca gaa ttc cga cat gac tca gga tat gaa gtt 98
     Ala Arg Ala Leu Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val
15
     15
                         20
                                             25
                                                                 30
     cat cat caa aaa ttg gtg ttc ttt gca gaa gat gtg ggt tca aac aaa
     His His Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
                      35
                                         40
                                                             45
20
     ggt gca atc att gga ctc atg gtg ggc ggt gtt gtc ata gcg act
                                                                    191
     Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Ile Ala Thr
                  50
                                      55
                                                         60
25
                                                                      197
     taagac
     <210>8
     <211> 61
30
     <212> PRT
     <213> Artificial Sequence
     <400>8
     Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala Trp Thr Ala Arg
35
       1
                       5
                                          10
                                                             15
```

Ala Leu Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His 25 20 30 Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala 5 35 40 45 Ile Ile Gly Leu Met Val Gly Gly Val Val Ile Ala Thr 50 55 60 10 <210> 9 <211> 137 <212> DNA <213> Artificial Sequence 15 <220> <223> Description of Artificial Sequence: DNA Construct <220> 20 <221> CDS <222> (17)..(133) <220> <221> misc_feature <222> (3)..(10) 25 <223> Not I recognition site <220> <221> misc_feature 30 <222> (11)..(16) <223> Xba I recognition site <220>

<221> sig_peptide

<222> (17)..(70)

```
<400>9
     tggcggccgc tctaga atg ctg ccc ggt ttg gca ctg ctc ctg ctg gcc gcc 52
                      Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala
                                        5
                                                          10
                        1
 5
     tgg acg gct cgg gcg ctt gat gca gaa ttc cga cat gac tca gga tat
     Trp Thr Ala Arg Ala Leu Asp Ala Glu Phe Arg His Asp Ser Gly Tyr
              15
                                 20
                                                     25
     gaa gtt cat cat caa aaa ttg gtg ttc ttt gct taag
10
                                                                  137
     Glu Val His His Gln Lys Leu Val Phe Phe Ala
          30
                              35
15
     <210> 10
     <211> 39
      <212> PRT
      <213> Artificial Sequence
     <400> 10
20
     Met Leu Pro Gly Leu Ala Leu Leu Leu Leu Ala Ala Trp Thr Ala Arg
       1
                       5
                                         10
                                                             15
     Ala Leu Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His
25
                  20
                                     25
                                                         30
     Gln Lys Leu Val Phe Phe Ala
              35
30
      <210> 11
      <211> 27
      <212> DNA
      <213> Artificial Sequence
35
```

```
<220>
     <223> Description of Artificial Sequence: Primer
     <400> 11
 5 gatgcagaat tccgacatga ctcagga
                                                                   27
     <210> 12
     <211> 28
10
     <212> DNA
     <213> Artificial Sequence
     <220>
     <223> Description of Artificial Sequence: Primer
15
     <400> 12
                                                                  28
     gtcttaagtc gctatgacaa caccgccc
20
     <210> 13
     <211> 62
     <212> DNA
     <213> Artificial Sequence
25
     <220>
     <223> Description of Artificial Sequence: Signal Sequence
           Adaptor
     <400> 13
     ggtctagaat gctgcccggt ttggcactgc tcctgctggc cgcctggacg gctcgggcgc 60
30
     tt
                                                                 62
     <210> 14
35
     <211> 61
     <212> DNA
```

	<213> Artificial Sequence	
	<220>	
	<223> Description of Artificial Sequence: Signal Sequence	
5	Adaptor	
	<400> 14	
	agcgcccgag ccgtccaggc ggccagcagg agcagtgcca aaccgggcag	cattctagac
10	С	61
	<210> 15	
	<211> 27	
15	<212> DNA	
13	<213> Artificial Sequence	
	<220>	
20	<223> Description of Artificial Sequence: Primer	
20	<400> 15	
	ggtctagaat gctgcccggt ttggcac	27
25	<210> 16	
	<211> 19	
	<212> DNA	
	<213> Artificial Sequence	
30	<220>	
	<223> Description of Artificial Sequence: Primer	
	<400> 16	
	tggcggccgc tctagaatg	19

	<210> 17	
	<211> 22	
	<212> DNA	
	<213> Artificial Sequence	
5		
	<220>	
	<223> Description of Artificial Sequence: Primer	
	<400> 17	
10	cacatcttaa gcaaagaaca cc	22
	<210> 18	
	<211> 18	
15	<212> DNA	
	<213> Artificial Sequence	
	<220>	
20	<223> Description of Artificial Sequence: Primer	
	<400> 18	
	agtgaaccgt cagatcgc	18
25	<210> 19	
	<211> 18	
	<212> DNA	
	<213> Artificial Sequence	
30	<220>	
	<223> Description of Artificial Sequence: Primer	
	<400> 19	
	cggtatcagc tcactcaa	18